> This Raw Listing contains the General Information
> Section and up to first 5 pages.

ENTERED

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 1   <110> APPLICANT: Fainzilber, Michael M.
 2         Kits, Karel S.
 3         Burlingame, Alma L.
 4         Olivera, Baldomero M.
 5         Walker, Craig
 6         Watkins, Maren
 7         Shetty, Reshma
 8         Cruz, Lourdes J.
 9         Imperial, Julita
10         Colledge, Clark
11         University of Utah Resarch Foundation
12         Vrije Universiteit
13         Regents of the University of California
14   <120> TITLE OF INVENTION: Gamma-Conopeptides
15   <130> FILE REFERENCE: Gamma-Conpeptides
16   <140> CURRENT APPLICATION NUMBER: US/09/210,952A
17   <141> CURRENT FILING DATE: 1998-12-15
18   <150> EARLIER APPLICATION NUMBER: US 60/069,706
19   <151> EARLIER FILING DATE: 1997-12-16
20   <160> NUMBER OF SEQ ID NOS: 47
21   <170> SOFTWARE: PatentIn Ver. 2.0
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23   <211> LENGTH: 42
24   <212> TYPE: PRT
25   <213> ORGANISM: Artificial Sequence
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27   <223> OTHER INFORMATION: Description of Artificial Sequence:generic formula
28         of gamma-conopeptides
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30   <221> NAME/KEY: PEPTIDE
31   <222> LOCATION: (1)..(13)
32   <223> OTHER INFORMATION: Xaa at residues 1, 2, 3, 4, 5, and 6 may be
33         des-Xaa or any amino acid; Xaa at residues 8, 9,
34         10, 11 and 12 may be any amino acid; Xaa at
35         residue 13 may be des-Xaa or any amino acid.
36   <220> FEATURE:
37   <221> NAME/KEY: PEPTIDE
38   <222> LOCATION: (15)..(19)
39   <223> OTHER INFORMATION: Xaa at residues 15, 16, 17 and 18 may be any amino
40         acid; Xaa at residue 19 is Glu,
41         gamma-carboxyglutamate or Gln.
42   <220> FEATURE:
43   <221> NAME/KEY: PEPTIDE
44   <222> LOCATION: (22)..(28)
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46          acid; Xaa at residue 25 may be des-Xaa or any
47          amino acid; Xaa at residues 27, 28 and 29 may be
48          any amino acid.
49    <220> FEATURE:
50    <221> NAME/KEY: PEPTIDE
51    <222> LOCATION: (30)..(42)
52    <223> OTHER INFORMATION: Xaa at residues 30, 31 and 32 may be des-Xaa or
53          any amino acid; Xaa at residues 34 and 35 may be
54          any amino acid; Xaa at residues 36, 37, 38, 39,
55          40, 41 and 42 may be des-Xaa or any  amino acid.
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68    <223> OTHER INFORMATION: Description of Artificial Sequence:generic
69          sequence of gamma-conopeptides.
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71    <221> NAME/KEY: PEPTIDE
72    <222> LOCATION: (1)..(13)
73    <223> OTHER INFORMATION: Xaa at residues 1, 2, 3, 4, 5 and 6 may be des-Xaa
74          or any amino acid; Xaa at residues 8, 9, 10, 11
75          and 12 may be any amino acid; Xaa at residue 13
76          may be des-Xaa or any amino acid.
77    <220> FEATURE:
78    <221> NAME/KEY: PEPTIDE
79    <222> LOCATION: (15)..(22)
80    <223> OTHER INFORMATION: Xaa at residues 15, 16, 17 and 18 may be any amino
81          acid; Xaa at residue 19 is Glu,
82          gamma-carboxyglutamate or Gln; Xaa at residue 22
83          is Ser or Thr.
84    <220> FEATURE:
85    <221> NAME/KEY: PEPTIDE
86    <222> LOCATION: (23)..(29)
87    <223> OTHER INFORMATION: Xaa at residues 23 and 24 may be any amino acid;
88          Xaa at residue 25 may be des-Xaa or any amino
89          acid; Xaa at residues 27, 28 and 29 may be any
90          amino acid.
91    <220> FEATURE:
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94    <223> OTHER INFORMATION: Xaa at residues 30, 31 and 32 may be des-Xaa or
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W-->

W-->

W-->

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       96          any amino acid; Xaa at residues 36, 37, 38, 39,
       97          40, 41 and 42 may be des-Xaa or any amino acid.
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       115   <223> OTHER INFORMATION: Xaa at residue 1 is any amino acid; Xaa at
       116          residues 2, 3, 4, 5 and 6 may be des-Xaa or any
       117          amino acid; Xaa at residues 8, 9, 10, 11, 12 and
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       124          and 39 may be des-Xaa or any amino acid.
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       130          gamma-carboxyglutamate or Gln.
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148  <223> OTHER INFORMATION: Xaa at residue 1 is any amino acid; Xaa at
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150        amino acid; Xaa at residues 8, 9, 10, 11, 12 and
151        13 may be any amino acid.
152  <220> FEATURE:
153  <221> NAME/KEY: PEPTIDE
154  <222> LOCATION: (15)..(19)
155  <223> OTHER INFORMATION: Xaa at residue 15 is Ser or Thr; Xaa at residues
156        16, 17 and 18 may ba any amino acid; Xaa at
157        residue 19 is Glu, gamma-carboxyglutamate or Gln.
158  <220> FEATURE:
159  <221> NAME/KEY: PEPTIDE
160  <222> LOCATION: (27)..(39)
161  <223> OTHER INFORMATION: Xaa at residues 27, 28, 29, 31, 32, 33, 34, 35, 36
162        and 37 may be any amino acid; Xaa at residues 38
163        and 39 may be des-Xaa or any amino acid.
164  <400> SEQUENCE: 4
165        Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa Xaa Xaa Xaa Xaa Cys Xaa Xaa
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167        Xaa Xaa Xaa Cys Cys Ser Asn Ser Cys Asp Xaa Xaa Xaa Cys Xaa Xaa
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173  <212> TYPE: PRT
174  <213> ORGANISM: Artificial Sequence
175  <220> FEATURE:
176  <223> OTHER INFORMATION: Description of Artificial Sequence:generic
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178  <220> FEATURE:
179  <221> NAME/KEY: PEPTIDE
180  <222> LOCATION: (1)..(6)
181  <223> OTHER INFORMATION: Xaa at residues 1 and 2 may be des-Xaa or a any
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183        gamma-carboxyglutamate; Xaa at residues 5 and 6
184        may be any amino acid.
185  <220> FEATURE:
186  <221> NAME/KEY: PEPTIDE
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188  <223> OTHER INFORMATION: Xaa at residue 7 is Trp or 6-bromo-Trp; Xaa at
189        residues 9, 10, 13 and 14 may be any amino acid;
190        Xaa at residue 16 is Glu, gamma-carboxyglutamate
191        or Gln.
192  <220> FEATURE:
193  <221> NAME/KEY: PEPTIDE
194  <222> LOCATION: (28)..(34)
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207    <220> FEATURE:
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209    <222> LOCATION: (1)..(31)
210    <223> OTHER INFORMATION: Xaa at residue 5 is Trp or 6-bromo-Trp; Xaa at
211          residues 14 and 26 are Glu or
212          gamma-carboxyglutamate; Xaa at residue 31 is Pro
213          or hydroxy-Pro.
214    <400> SEQUENCE: 6
215          Asp Cys Thr Ser Xaa Phe Gly Arg Cys Thr Val Asn Ser Xaa Cys Cys
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222    <213> ORGANISM: Conus textile
223    <220> FEATURE:
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225    <222> LOCATION: (1)..(34)
226    <223> OTHER INFORMATION: Xaa at residues 1, 7 and 34 are Trp or
227          6-bromo-Trp; Xaa at residues 3 and 16 are Glu or
228          gamma-carboxyglutamate; Xaa at residues 31 and 32
229          are Pro or hydroxy-Pro.
230    <400> SEQUENCE: 7
231          Xaa Leu Xaa Cys Ser Val Xaa Phe Ser His Cys Thr Lys Asp Ser Xaa
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233          Cys Cys Ser Asn Ser Cys Asp Gln Thr Tyr Cys Thr Leu Met Xaa Xaa
234                       20              25              30
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236    <210> SEQ ID NO 8
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239    <213> ORGANISM: Conus textile
240    <220> FEATURE:
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242    <222> LOCATION: (1)..(39)
243    <223> OTHER INFORMATION: Xaa at residues 1, 2, 4, 10 and 39 are Trp or
                6-bromo-Trp ; Xaa at residues 19 and 31 are Glu or
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Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220> to
<223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY
PATENT APPLICATION **US/09/210,952A**

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